

Sequence Listing.ST25.txt
SEQUENCE LISTING

<110> Pfizer Ltd. (EP (GB) only)
Pfizer Inc. (US, JP, EP except GB)
Fidock, Mark David

<120> Novel Polypeptide

<130> PC10960AGPR

<150> GB 0030855.1

<151> 2000-12-18

<150> GB 0101222.8

<151> 2001-01-17

<160> 7

<170> PatentIn version 3.0

<210> 1

<211> 1082

<212> DNA

<213> Homo sapiens

<400> 1

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ggtctctgcc agttctcaga gaagtacaag caagtctacc tctccctggc ctacagtatc	180
atcttttatcc tagggctgcc actaaatggc actgtcttgt ggcaactcctg gggccaaacc	240
aagcgctgga gctgtgccac cacctatctg gtgaacctga tggtagccga cctgctttat	300
gtgctattgc ccttcctcat catcacctac tcactagatg acaggtggcc cttcggggag	360
ctgctctgca agctgggtgca cttcctgttc tatatcaacc ttacggcag catcctgctg	420
ctgacctgca tctctgtgca ccagttccta ggtgtgtgcc accactgtg ttcgctgccc	480
taccggaccc gcaggcatgc ctggctgggc accagcacca cctgggccct ggtggtcctc	540
cagctgctgc ccacactggc cttctccac acggactaca tcaatggcca gatgatctgg	600
tatgacatga ccagccaaga gaattttgat cggctttttg cctacggcat agttctgaca	660
ttgtctggct ttctttccct ccttggctcat ttggtgtgc tattcactga tggtcaggag	720
cctgatcaag ccagaggaga acctcatgag gacaggcaac acagcccgag ccaggtccat	780
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tcgctccttc tacctcacca tctgctttct gctttctcag gactgccagc tcttgatggc      900
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agtcctgtac tttctttcaa ggggggcaaa aatagagtca ggctcctcca gaaactgagg    1020
cagaacaagt tgggtgagca tccagctggg aggaagagat gccaggggtt gaacagatct    1080
gg                                     1082

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<210> 2
<211> 360
<212> PRT
<213> Homo sapiens

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<400> 2

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Met Leu Ser Ile Leu Leu Pro Ser Arg Gly Ser Arg Ser Gly Ser Arg
1          5          10          15
Arg Gly Ala Leu Leu Leu Glu Gly Ala Ser Arg Asp Met Glu Lys Val
20          25          30
Asp Met Asn Thr Ser Gln Glu Gln Gly Leu Cys Gln Phe Ser Glu Lys
35          40          45
Tyr Lys Gln Val Tyr Leu Ser Leu Ala Tyr Ser Ile Ile Phe Ile Leu
50          55          60
Gly Leu Pro Leu Asn Gly Thr Val Leu Trp His Ser Trp Gly Gln Thr
65          70          75          80
Lys Arg Trp Ser Cys Ala Thr Thr Tyr Leu Val Asn Leu Met Val Ala
85          90          95
Asp Leu Leu Tyr Val Leu Leu Pro Phe Leu Ile Ile Thr Tyr Ser Leu
100         105         110
Asp Asp Arg Trp Pro Phe Gly Glu Leu Leu Cys Lys Leu Val His Phe
115         120         125
Leu Phe Tyr Ile Asn Leu Tyr Gly Ser Ile Leu Leu Leu Thr Cys Ile
130         135         140
Ser Val His Gln Phe Leu Gly Val Cys His Pro Leu Cys Ser Leu Pro
145         150         155         160
Tyr Arg Thr Arg Arg His Ala Trp Leu Gly Thr Ser Thr Thr Trp Ala
165         170         175
Leu Val Val Leu Gln Leu Leu Pro Thr Leu Ala Phe Ser His Thr Asp

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180

185

190

Tyr Ile Asn Gly Gln Met Ile Trp Tyr Asp Met Thr Ser Gln Glu Asn
 195 200 205

Phe Asp Arg Leu Phe Ala Tyr Gly Ile Val Leu Thr Leu Ser Gly Phe
 210 215 220

Leu Ser Leu Leu Gly His Phe Gly Val Leu Phe Thr Asp Gly Gln Glu
 225 230 235 240

Pro Asp Gln Ala Arg Gly Glu Pro His Glu Asp Arg Gln His Ser Pro
 245 250 255

Ser Gln Val His Pro Asp His Pro Thr Gly Val Trp Pro Leu His Pro
 260 265 270

Leu Phe Cys Ala Leu Pro Tyr His Ser Leu Leu Leu Pro His His Leu
 275 280 285

Leu Ser Ala Phe Ser Gly Leu Pro Ala Leu Asp Gly Ser Gln Cys Gly
 290 295 300

Leu Gln Asp Met Glu Ala Ser Gly Glu Cys Glu Gln Leu Pro Gln Pro
 305 310 315 320

Ser Pro Val Leu Ser Phe Lys Gly Gly Lys Asn Arg Val Arg Leu Leu
 325 330 335

Gln Lys Leu Arg Gln Asn Lys Leu Gly Glu His Pro Ala Gly Arg Lys
 340 345 350

Arg Cys Pro Gly Leu Asn Arg Ser
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<210> 3

<211> 1020

<212> DNA

<213> Homo sapiens

<400> 3

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ggctctctgcc agttctcaga gaagtacaag caagtctacc tctccctggc ctacagtatc 180

atcttttatcc tagggctgcc actaaatggc actgtcttgt ggcactcctg gggccaaacc 240

aagcgctgga gctgtgccac cacctatctg gtgaacctga tgggtggccga cctgctttat 300

gtgctattgc ccttcctcat catcacctac tcactagatg acaggtggcc cttcggggag 360

Sequence Listing.ST25.txt

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ctgctctgca agctggtgca cttcctgttc tatatcaacc ttacggcag catcctgctg 420
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tacgggaccc gcaggcatgc ctggctgggc accagcacca cctgggacct ggtggtctc 540
cagctgctgc ccacactggc cttctccac acggactaca tcaatggcca gatgatctgg 600
tatgacatga ccagccaaga gaattttgat cggctttttg cctacggcat agttctgaca 660
ttgtctggct ttctttcccc ctcttggtc attttggtgt gctattcact gatggtcagg 720
agcctgatca agccagagga gaacctcatg aggacaggca acacagcccg agccaggctc 780
atccggacca tcctactggt gtgtggcctc ttcacctct gttttgtgcc ctccatatac 840
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gcaccacgtg tggcctacaa gatatggagg cctctggtga gtgtgagcag ctgcctcaac 960
ccagtcctgt actttctttc aaggggggca aaaatagagt caggctcctc cagaaactga 1020

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<210> 4
<211> 339
<212> PRT
<213> Homo sapiens

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<400> 4

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```

Met Leu Ser Ile Leu Leu Pro Ser Arg Gly Ser Arg Ser Gly Ser Arg
1      5      10      15
Arg Gly Ala Leu Leu Leu Glu Gly Ala Ser Arg Asp Met Glu Lys Val
20      25      30
Asp Met Asn Thr Ser Gln Glu Gln Gly Leu Cys Gln Phe Ser Glu Lys
35      40      45
Tyr Lys Gln Val Tyr Leu Ser Leu Ala Tyr Ser Ile Ile Phe Ile Leu
50      55      60
Gly Leu Pro Leu Asn Gly Thr Val Leu Trp His Ser Trp Gly Gln Thr
65      70      75      80
Lys Arg Trp Ser Cys Ala Thr Thr Tyr Leu Val Asn Leu Met Val Ala
85      90      95
Asp Leu Leu Tyr Val Leu Leu Pro Phe Leu Ile Ile Thr Tyr Ser Leu
100     105     110
Asp Asp Arg Trp Pro Phe Gly Glu Leu Leu Cys Lys Leu Val His Phe
115     120     125

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Sequence Listing.ST25.txt

```

Leu Phe Tyr Ile Asn Leu Tyr Gly Ser Ile Leu Leu Leu Thr Cys Ile
 130      135      140

Ser Val His Gln Phe Leu Gly Val Trp His Pro Leu Cys Ser Leu Pro
 145      150      155      160

Tyr Arg Thr Arg Arg His Ala Trp Leu Gly Thr Ser Thr Thr Trp Ala
      165      170      175

Leu Val Val Leu Gln Leu Leu Pro Thr Leu Ala Phe Ser His Thr Asp
      180      185      190

Tyr Ile Asn Gly Gln Met Ile Trp Tyr Asp Met Thr Ser Gln Glu Asn
      195      200      205

Phe Asp Arg Leu Phe Ala Tyr Gly Ile Val Leu Thr Leu Ser Gly Phe
      210      215      220

Leu Ser Pro Ser Leu Val Ile Leu Val Cys Tyr Ser Leu Met Val Arg
      225      230      235      240

Ser Leu Ile Lys Pro Glu Glu Asn Leu Met Arg Thr Gly Asn Thr Ala
      245      250      255

Arg Ala Arg Ser Ile Arg Thr Ile Leu Leu Val Cys Gly Leu Phe Thr
      260      265      270

Leu Cys Phe Val Pro Phe His Ile Thr Arg Ser Phe Tyr Leu Thr Ile
      275      280      285

Cys Phe Leu Leu Ser Gln Asp Cys Gln Leu Leu Met Ala Pro Ser Val
      290      295      300

Ala Tyr Lys Ile Trp Arg Pro Leu Val Ser Val Ser Ser Cys Leu Asn
      305      310      315      320

Pro Val Leu Tyr Phe Leu Ser Arg Gly Ala Lys Ile Glu Ser Gly Ser
      325      330      335

Ser Arg Asn

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```

<210> 5
<211> 27
<212> DNA
<213> Homo sapiens

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<400> 5
accatgctgt ccattttgct tccttcc

```

27

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<210> 6

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Sequence Listing.ST25.txt

<211> 24
<212> DNA
<213> Homo sapiens

<400> 6
tcaccagatc tgttcaaccc tggg 24

<210> 7
<211> 24
<212> DNA
<213> Homo sapiens

<400> 7
tcagtttctg gaggagcctg actc 24